



2570  
1001

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/052,648  
Source: OIPE  
Date Processed by STIC: 10/1/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

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<http://www.uspto.gov/web/offices/pac/checker>

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
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3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
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U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202



OIPE

## RAW SEQUENCE LISTING

DATE: 10/01/2002

PATENT APPLICATION: US/10/052,648

TIME: 16:42:28

Input Set : A:\Cura5501.app

Output Set: N:\CRF4\10012002\J052648.raw

3 <110> APPLICANT: Anderson, David  
4 Burgess, Catherine  
5 Casman, Stacie  
6 Colman, Steven  
7 Edinger, Shlomit R.  
8 Ellerman, Karen  
9 Gerlach, Valerie  
10 Gunther, Erik  
11 Kekuda, Ramesh  
12 MacDougall, John R.  
13 Mehraban, Fuad  
14 Patturajan, Meera  
15 Rothenberg, Mark  
16 Shimkets, Richard  
17 Smithson, Glennnda  
18 Spytek, Kimberly A.  
19 Stone, David J.  
20 Vernet, Corine A.M.  
21 Zerhusen, Bryan D.  
23 <120> TITLE OF INVENTION: PROTEINS, POLYNUCLEOTIDES ENCODING THEM AND METHODS OF  
24 USING THE SAME  
26 <130> FILE REFERENCE: 21402-250 (CURA-550)  
28 <140> CURRENT APPLICATION NUMBER: 10/052,648  
29 <141> CURRENT FILING DATE: 2002-01-18  
31 <150> PRIOR APPLICATION NUMBER: 60/262,454  
32 <151> PRIOR FILING DATE: 2001-01-18  
34 <150> PRIOR APPLICATION NUMBER: 60/272,920  
35 <151> PRIOR FILING DATE: 2001-03-02  
37 <150> PRIOR APPLICATION NUMBER: 60/284,549  
38 <151> PRIOR FILING DATE: 2001-04-18  
40 <150> PRIOR APPLICATION NUMBER: 60/303,229  
41 <151> PRIOR FILING DATE: 2001-07-05  
43 <150> PRIOR APPLICATION NUMBER: 60/262,892  
44 <151> PRIOR FILING DATE: 2001-01-19  
46 <150> PRIOR APPLICATION NUMBER: 60/263,605  
47 <151> PRIOR FILING DATE: 2001-01-23  
49 <150> PRIOR APPLICATION NUMBER: 60/269,098  
50 <151> PRIOR FILING DATE: 2001-02-15  
52 <150> PRIOR APPLICATION NUMBER: 60/264,159  
53 <151> PRIOR FILING DATE: 2001-01-25  
55 <150> PRIOR APPLICATION NUMBER: 60/265,517  
56 <151> PRIOR FILING DATE: 2001-01-31  
58 <150> PRIOR APPLICATION NUMBER: 60/271,855

Does Not Comply  
Corrected Diskette Needed  
PP. 2-4-5

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59 <151> PRIOR FILING DATE: 2001-02-27  
 61 <150> PRIOR APPLICATION NUMBER: 60/267,057  
 62 <151> PRIOR FILING DATE: 2001-02-07  
 64 <150> PRIOR APPLICATION NUMBER: 60/286,287  
 65 <151> PRIOR FILING DATE: 2001-04-25  
 67 <160> NUMBER OF SEQ ID NOS: 97  
 69 <170> SOFTWARE: PatentIn Ver. 2.1

## ERRORED SEQUENCES

3711 <210> SEQ ID NO: 35  
 3712 <211> LENGTH: 969  
 3713 <212> TYPE: PRT  
 3714 <213> ORGANISM: Homo sapiens  
 3716 <400> SEQUENCE: 35

*see p. 4*

3717	Met	His	Thr	Pro	Ser	Ile	Arg	Ser	Ile	Thr	His	Asp	Ala	Gln	Thr	Ser
3718	1				5					10					15	
3720	Ser	Thr	Gly	Ser	Ser	Ala	Pro	Gly	Thr	Ala	Leu	Cys	Thr	Glu	Glu	Cys
3721				20					25					30		
3723	Val	His	Gly	Arg	Cys	Val	Ser	Pro	Asp	Thr	Cys	His	Cys	Glu	Pro	Gly
3724			35					40					45			
3726	Trp	Gly	Gly	Pro	Asp	Cys	Ser	Ser	Gly	Cys	Asp	Ser	Asp	His	Trp	Gly
3727		50					55				60					
3729	Pro	His	Cys	Ser	Asn	Arg	Cys	Gln	Cys	Gln	Asn	Gly	Ala	Leu	Cys	Asn
3730	65				70					75					80	
3732	Pro	Ile	Thr	Gly	Ala	Cys	Val	Cys	Ala	Ala	Gly	Phe	Arg	Gly	Trp	Arg
3733				85						90					95	
3735	Cys	Glu	Glu	Leu	Cys	Ala	Pro	Gly	Thr	His	Gly	Lys	Gly	Cys	Gln	Leu
3736				100					105					110		
3738	Pro	Cys	Gln	Cys	Arg	His	Gly	Ala	Ser	Cys	Asp	Pro	Arg	Ala	Gly	Glu
3739			115				120						125			
3741	Cys	Leu	Cys	Ala	Pro	Gly	Tyr	Thr	Gly	Val	Tyr	Cys	Glu	Glu	Leu	Cys
3742		130					135					140				
3744	Pro	Pro	Gly	Ser	His	Gly	Ala	His	Cys	Glu	Leu	Arg	Cys	Pro	Cys	Gln
3745	145				150					155					160	
3747	Asn	Gly	Gly	Thr	Cys	His	His	Ile	Thr	Gly	Glu	Cys	Ala	Cys	Pro	Pro
3748				165						170					175	
3750	Gly	Trp	Thr	Gly	Ala	Val	Cys	Ala	Gln	Pro	Cys	Pro	Pro	Gly	Thr	Phe
3751				180					185					190		
3753	Gly	Gln	Asn	Cys	Ser	Gln	Asp	Cys	Pro	Cys	His	His	Gly	Gly	Gln	Cys
3754			195					200					205			
3756	Asp	His	Val	Thr	Gly	Gln	Cys	His	Cys	Thr	Ala	Gly	Tyr	Met	Gly	Asp
3757		210					215					220				
3759	Arg	Cys	Gln	Glu	Glu	Cys	Pro	Phe	Gly	Ser	Phe	Gly	Phe	Gln	Cys	Ser
3760	225					230					235				240	
3762	Gln	His	Cys	Asp	Cys	His	Asn	Gly	Gly	Gln	Cys	Ser	Pro	Thr	Thr	Gly
3763				245					250					255		
3765	Ala	Cys	Glu	Cys	Glu	Pro	Gly	Tyr	Lys	Gly	Pro	Arg	Cys	Gln	Glu	Arg

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3766		260		265		270
3768	Leu Cys Pro	Glu Gly Leu His Gly	Pro Gly Cys Thr	Leu Pro Cys Pro		
3769		275		280		285
3771	Cys Asp Ala	Asp Asn Thr Ile Ser	Cys His Pro Val	Thr Gly Ala Cys		
3772		290		295		300
3774	Thr Cys Gln	Pro Gly Trp Ser Gly	His His Cys Asn	Glu Ser Cys Pro		
3775	305		310		315	320
3777	Val Gly Tyr	Tyr Gly Asp Gly	Cys Gln Leu Pro	Cys Thr Cys Gln	Asn	
3778		325		330		335
3780	Gly Ala Asp	Cys His Ser Ile Thr	Gly Gly Cys Thr	Cys Ala Pro Gly		
3781		340		345		350
3783	Phe Met Gly	Glu Val Cys Ala Val	Ser Cys Ala Ala	Gly Thr Tyr Gly		
3784		355		360		365
3786	Pro Asn Cys	Ser Ser Ile Cys Ser	Cys Asn Asn Gly	Gly Thr Cys Ser		
3787		370		375		380
3789	Pro Val Asp	Gly Ser Cys Thr	Cys Lys Glu Gly	Trp Gln Gly Leu	Asp	
3790	385		390		395	400
3792	Cys Thr Leu	Pro Cys Pro Ser	Gly Thr Trp Gly	Leu Asn Cys Asn	Glu	
3793		405		410		415
3795	Ser Cys Thr	Cys Ala Asn Gly	Ala Ala Cys Ser	Pro Ile Asp Gly	Ser	
3796		420		425		430
3798	Cys Ser Cys	Thr Pro Gly Trp	Leu Gly Asp Thr	Cys Glu Leu Pro	Cys	
3799		435		440		445
3801	Pro Asp Gly	Thr Phe Gly Leu	Asn Cys Ser Glu	His Cys Asp Cys	Ser	
3802		450		455		460
3804	His Ala Asp	Gly Cys Asp Pro	Val Thr Gly His	Cys Cys Cys Leu	Ala	
3805	465		470		475	480
3807	Gly Trp Thr	Gly Ile Arg Cys	Asp Ser Thr Cys	Pro Pro Gly Arg	Trp	
3808		485		490		495
3810	Gly Pro Asn	Cys Ser Val Ser	Cys Ser Cys Glu	Asn Gly Gly Ser	Cys	
3811		500		505		510
3813	Ser Pro Glu	Asp Gly Ser Cys	Glu Cys Ala Pro	Gly Phe Arg Gly	Pro	
3814		515		520		525
3816	Leu Cys Gln	Arg Ile Cys Pro	Gly Phe Tyr Gly	His Gly Cys Ala		
3817		530		535		540
3819	Gln Pro Cys	Pro Leu Cys Val	His Ser Ser Arg	Pro Cys His His	Ile	
3820	545		550		555	560
3822	Ser Gly Ile	Cys Glu Cys Leu	Pro Gly Phe Ser	Gly Ala Leu Cys	Asn	
3823		565		570		575
3825	Gln Val Cys	Ala Gly Gly Tyr	Phe Gly Gln Asp	Cys Ala Gln Leu	Cys	
3826		580		585		590
3828	Ser Cys Ala	Asn Asn Gly Thr	Cys Ser Pro Ile	Asp Gly Ser Cys	Gln	
3829		595		600		605
3831	Cys Phe Pro	Gly Trp Ile Gly	Lys Asp Cys Ser	Gln Ala Cys Pro	Pro	
3832		610		615		620
3834	Gly Phe Trp	Gly Pro Ala Cys	Phe His Ala Cys	Ser Cys His Asn	Gly	
3835	625		630		635	640
3837	Ala Ser Cys	Ser Ala Glu Asp	Gly Ala Cys His	Cys Thr Pro Gly	Trp	
3838		645		650		655

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Input Set : A:\Cura5501.app

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```

3840 Thr Gly Leu Phe Cys Thr Gln Arg Cys Pro Ala Ala Phe Phe Gly Lys
3841                660                665                670
3843 Asp Cys Gly Arg Val Cys Gln Cys Gln Asn Gly Ala Ser Cys Asp His
3844                675                680                685
3846 Ile Ser Gly Lys Cys Thr Cys Arg Thr Gly Phe Thr Gly Gln His Cys
3847                690                695                700
3849 Glu Gln Arg Cys Ala Pro Gly Thr Phe Gly Tyr Gly Cys Gln Gln Leu
3850 705                710                715                720
3852 Cys Glu Cys Met Asn Asn Ser Thr Cys Asp His Val Thr Gly Thr Cys
3853                725                730                735
3855 Tyr Cys Ser Pro Gly Phe Lys Gly Ile Arg Cys Asp Gln Ala Ala Leu
3856                740                745                750
3858 Met Met Glu Glu Leu Asn Pro Tyr Thr Lys Ile Ser Pro Ala Leu Gly
3859                755                760                765
3861 Ala Glu Arg His Ser Val Gly Ala Val Thr Gly Ile Met Leu Leu Leu
3862                770                775                780
3864 Phe Leu Ile Val Val Leu Leu Gly Leu Phe Ala Trp His Arg Arg Arg
3865 785                790                795                800
3867 Gln Lys Glu Lys Gly Arg Asp Leu Ala Pro Arg Val Ser Tyr Thr Pro
3868                805                810                815
3870 Ala Met Arg Met Thr Ser Thr Asp Tyr Ser Leu Ser Gly Ala Cys Gly
3871                820                825                830
E--> 3873 Met Asp Arg Arg Gln Asn Thr Tyr Ile Met Asp Lys Gly Phe Lys Xaa
3874                835                840                845
E--> 3876 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
3877                850                855                860
E--> 3879 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
3880 865                870                875                880
E--> 3882 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Met Lys Ser Pro Val His Met
3883                885                890                895
3885 Gly Ser Pro Tyr Thr Asp Val Pro Ser Leu Ser Thr Ser Asn Lys Asn
3886                900                905                910
3888 Ile Tyr Glu Val Glu Pro Thr Val Ser Val Val Gln Glu Gly Cys Gly
3889                915                920                925
3891 His Asn Ser Ser Tyr Ile Gln Asn Ala Tyr Asp Leu Pro Arg Asn Ser
3892                930                935                940
3894 His Ile Pro Gly His Tyr Asp Leu Leu Pro Val Arg Gln Ser Pro Ala
3895 945                950                955                960
3897 Asn Gly Pro Ser Gln Asp Lys Gln Ser
3898                965

```

*see p.5  
for error  
explanation*

## VARIABLE LOCATION SUMMARY

DATE: 10/01/2002

PATENT APPLICATION: US/10/052,648

TIME: 16:42:30

Input Set : A:\Cura5501.app

Output Set: N:\CRF4\10012002\J052648.raw

Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of &lt;220&gt; to &lt;223&gt; is MANDATORY if n's or Xaa's are present.

in &lt;220&gt; to &lt;223&gt; section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:13; N Pos. 1408

Seq#:14; Xaa Pos. 203

Seq#:15; N Pos. 2004

Seq#:35; Xaa Pos. 848,849,850,851,852,853,854,855,856,857,858,859,860,861

Seq#:35; Xaa Pos. 862,863,864,865,866,867,868,869,870,871,872,873,874,875

Seq#:35; Xaa Pos. 876,877,878,879,880,881,882,883,884,885,886,887,888,889

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/052,648

DATE: 10/01/2002

TIME: 16:42:30

Input Set : A:\Cura5501.app

Output Set: N:\CRF4\10012002\J052648.raw

L:1553 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1380  
L:1624 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:192  
L:1696 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1980  
L:3873 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:35  
M:340 Repeated in SeqNo=35